

SEQ ID NO: 11 CCR5 F P Y S Q Y Q P W K N F Q T L K I V I L G L V L P
 SEQ ID NO: 12 TTCCATACAGtcagtcagtaatcaattcttgaagaattccagacaTAAAGATAGTCATCTTGGGCTGGTCTCTGCCG
 SEQ ID NO: 13 Δccr5 P P Y I K D S H L G A G P A

deletion

CCR5 L L V M V I C Y S G I L K T L L R C R N E K K R
CTGCTTGTCATGGTCATCTGCTACTCGGAATCCTAAAACTCTGCTTCGGTGTCCAATGAGAAGAGAGG
 Δccr5 A A C H G H L L L G N P K N S A S V S K *

FIG. 6B

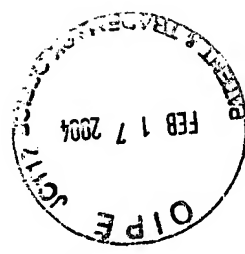


FIG. 2A
FIG. 2B

FIG. 2

SEQ ID NO: 5	CCR5	1	MDYQVSSPH	DINVTSEPCQKINVKQIMAK	LLPPLYSLVIFGFGVGNMLVIL	ILINCKRKLSMTDIYLLNLAISDILFL	83	
SEQ ID NO: 7	hcc-R2b	6	MLSTSRSRFRINWNEEGEVTFFD	YDYGAPCHKFDVKQISAG	LLPPLYSLVIFGFGVGNMLVIL	ILINCKRKLSMTDIYLLNLAISDILFL	95	
SEQ ID NO: 8	hcc-R3		MTTSIDTVEFTGTSYYDDVGLIQEKADTRALMAQFV	PPLYSLVFTVGLIGNVVMILIKYRIRI	MTNIIYLLNLAISDILFL		87	
SEQ ID NO: 9	hcc-R1		METPNATEDYDTTEFD	IGDAPCQKINERAFGALL	PPLYSLVFVIGVGNMLVIL	VQVQKRLKMTSIYLLNLAISDILFL	87	
SEQ ID NO: 10	hcc-R4		NNPTDIADTTIDESIYSNVLYESIPKPTKEGKAFGELF	LPPLYSLVFVIGVGNMLVIL	VFKYKRIRSMTDVYLLNLAISDILFL		92	
II								
SEQ ID NO: 5	CCR5		VPFWAHYA	ADQWDEGNINCOLLTGLYF	IGFTSGIFFIILLTIDRYLA	WHAVALKARTVTFGVVTSVITW	VAVFASLPGIIFTRSQEGIH	177
SEQ ID NO: 7	hcc-R2b		LPLWAHSA	ANENWFGNACKLFTGLYHI	IGYEGSIFFIILLTIDRYLA	IVHAVFALKARTVTFGVVTSVITW	VAVFASVPGIIFTKQKEDSV	189
SEQ ID NO: 8	hcc-R3		LPFWIHYVRGHVWFGHGNLISGFVHTGLYSEL	FFIILLTIDRYLA	IVHAVFALKARTVTFGVVTSVITW	WGLAVHAAALPEFI	FYETEELFEE	182
SEQ ID NO: 9	hcc-R1		LPFWIDYKLKDDAVFGDAMCKILISFWTGLYSEL	FFIILLTIDRYLA	IVHAVFALKARTVTFGVVTSVITW	ALALIASMPGLYFSKTQ	WEFTIH	182
SEQ ID NO: 10	hcc-R4		LPFWGYIA	ADQWVFGLGICKMISWMLVGFYSGIFFVFMMSIDRYLA	IVHAVFSLRARTL	IVGVITSLATWSVAVFASLPGIIFTRSQEGIH		186
IV								
III								

FIG. 2A